Applicant: Sunghwa Choe, et al.

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## In the specification:

Please amend the paragraph beginning at page 18, line 17 to page 19, line 8, line as follows:

The term encompasses alterations to the polynucleotide sequence, so long as the alteration results in a plant displaying one or more dwf7 phenotypic traits (described below) when the polynucleotide is expressed in a plant. Such modifications typically include deletions, additions, and substitutions, to the native dwf7 sequence, so long as the mutation results in a plant displaying a dwf7 phenotype as defined below. These modifications may be deliberate, as through site-directed mutagenesis, or may be accidental, such as through mutations of plants which express the dwf7 polynucleotide or errors due to PCR amplification. The term encompasses expressed allelic variants of the wild-type dwf7 sequence which may occur by normal genetic variation or are produced by genetic engineering methods and which result in a detectable change in the wild-type dwf7 phenotype. Two particular dwf7 allelic variants described herein are dwf7-1 and dwf7-2. Polypeptides corresponding to these variants include about amino acids 1-60 1-230 and 1-230 1-60, respectively, of Figure 9. However, the boundaries of these polypeptides may vary by 1 to 10 or more amino acids, or any integer therebetween. Thus, dwf7-1 and dwf7-2 polypeptides may include, for example, amino acids 1-229  $\frac{1-59}{1-59}$  and  $\frac{1-59}{1-229}$ , respectively, or  $\frac{3-232}{1-29}$  and  $\frac{3-62}{1-29}$ , respectively, and so on. Also described herein is a dwf7 polynucleotide termed "HDF7." The term "dwf7 polynucleotide" as used herein, is intended to encompass the HDF7 polynucleotide. This polynucleotide is shown in Figures 10A-10F herein. The polypeptide encoded by HDF7 is depicted at about positions 1-279 of Figure 11. As with the dwf7-1 and dwf7-2 polypeptides, the boundaries of the HDF7 polypeptide may also vary by 1 to 10 or more amino acids, or any integer therebetween. These molecules are discussed in detail below.